

## (1) GENERAL INFORMATION:

- (i) APPLICANTS: HU, A-LI W. HARTLEY, JAMES L. JORDAN, HEATHER J.
- (ii) TITLE OF INVENTION: NUCLEIC ACID LADDERS
- (iii) NUMBER OF SEQUENCES: 7
  - (iv) CORRESPONDENCE ADDRESS:
    - (A) ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX, P.L.L.C.

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- (B) STREET: 1100 NEW YORK AVENUE N.W., SUITE 600
- (C) CITY: WASHINGTON
- (D) STATE: DC
- (E) COUNTRY: USA
- (F) ZIP: 20005
- (v) COMPUTER READABLE FORM:
  - (A) MEDIUM TYPE: Floppy disk
  - (B) COMPUTER: IBM PC compatible
  - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
  - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:
  - (A) APPLICATION NUMBER: (To be assigned)
  - (B) FILING DATE: (Herewith)
  - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
  - (A) APPLICATION NUMBER: US 09/048,242
  - (B) FILING DATE: 26-MAR-1998
  - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
  - (A) APPLICATION NUMBER: US 60/040,914
  - (B) FILING DATE: 27-MAR-1997
  - (C) CLASSIFICATION:
- (ix) TELECOMMUNICATION INFORMATION:
  - (A) TELEPHONE: (202)371-2600
  - (B) TELEFAX: (202)371-2540
- (2) INFORMATION FOR SEQ ID NO:1:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 10 base pairs
    - (B) TYPE: nucleic acid
    - (C) STRANDEDNESS: both
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: cDNA
  - (x1) SEQUENCE DESCRIPTION: SEQ ID NO:1:

ATC	CAGGAT	10
(2)	INFORMATION FOR SEQ ID NO:2:	
	<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 10 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: both</li><li>(D) TOPOLOGY: linear</li></ul>	
	(ii) MOLECULE TYPE: cDNA	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
ATC	AGTCGAT .	10
(2)	INFORMATION FOR SEQ ID NO:3:	
	(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 10 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: both  (D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: cDNA	
	/wil SEQUENCE DESCRIPTION, SEC ID NO.3.	
<u>አ</u> መርረ	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	1.0
	TATEORNATION FOR GEO. ID. NO. 4:	10
(2)	INFORMATION FOR SEQ ID NO:4:	
	<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 10 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: both</li><li>(D) TOPOLOGY: linear</li></ul>	
•	(ii) MOLECULE TYPE: cDNA	
	/wil Chounge Decontration, Cho ID NO. 4	
7 m ~ :	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	1.0
	ATGCGAT	10
(2)	INFORMATION FOR SEQ ID NO:5:	
	<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 65 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: both</li><li>(D) TOPOLOGY: linear</li></ul>	
	(ii) MOLECULE TYPE: cDNA	

(	xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
AATTC	TCGGG ATATCTCAGG ATATCTCAGG ATATCTCAGG ATATCTCGGG	60
CTGCA		65
(2) I	NFORMATION FOR SEQ ID NO:6:	
	(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 43 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(	ii) MOLECULE TYPE: cDNA	
(	xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
CAACA	ARCAAG AATTCAGGCC TAGCCAGTGC CTCGTCCATT TTT	43
(2) I	NFORMATION FOR SEQ ID NO:7:	
	(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 37 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(	(ii) MOLECULE TYPE: cDNA	
(	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
CAACA	AACAAA AGCTTCTTAC ATGGCCCAGG TGCAGTA	37